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                                                                                                                                                                                                                                                                                                                                                                                                           AF125672 8686 bp mRNA linear PRI 04-AF. Homo sapiens silencing mediator of retinoic acid and thyroid hormone receptor extended isoform (SMRTE) mRNA, complete cds. AF125672
                                                                                                                                                                                                                                 Park, B.J., Schroen, D.J., Yang, M., Li, H., Li, L. and Chen, J.D. SMRTe, a silencing mediator for retinoid and thyroid hormone receptors-extended isoform that is more related to the nuclear
                                                                                         Submitted (03-FEB-1999) Pharmacology and Molecular Toxicology, University of Massachusetts Medical School, 55 Lake Avenue North, Worcester, MA 01655, USA
                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                        Chen, J.D.
Direct Submission
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/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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S 밁 S 용 Ş 밁 Query Match Best Local Similarity Matches 8472; 181 156 13 GCCCCAGCGGCGGAAGGCCCTCCCTGCTGTCTGAGTTCCAGCCCGGGAATGAACGGTCCCA GGAGTACCAGCACCACTCCCGCGACTATGCCTCCCACCTGTCGCCGGGCTCCATCATCCA 180 GCCCCACAGCCTTTCCTACCCAGTGCAGATCGCCCGGACGCACACGGACGTCGGGCTCCT 120 CATGTCGGGCTCCACACAGCCTGTGGCACAGACGTGGAGGGCCACTGAGCCCCGCTACCC CATGTCGGGCTCCACACACACGCTTGTGGCACAGACGTGGAGGGCCACTGAGCCCCGGCTACCCC GCCCCACAGCCTTTCCTACCCAGTGCAGATCGCCCGGACGCACACGGACGTCGGGCTCCT Conservative 97.2%; Score 8324.6; Pred. No. 0; Mismatches B 34; 8 Indels Length 8686; 88; Gaps 240 335 275 215 60

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2382 GHACHAGHGHANAANGGGCCCAAGCCCCAGCCACCCCCCCCGGGCCCCCCCC		2161 GGCTGAAGCCTTACATGCCTCTGGGAATGAGGTGCCCAGAGGGGAATGCAGTGGCCCAGC 2220 2313 GGCTGAAGC 2321 2313 GGCTGACAACAGCTCAGACACCGAGAGAGCATCCCCTCTCCTCACACTGAGGCCGCCAA 2280	GGAGG	041 193	1981 GAACCTCGATGAGATCTTGCAGCAGCAGACTGAAGATGGAGAAGGAGGAGGAACGCGCGC 2040 	21 73	1861 GGAAACAGCCAAGAAAGGTCTCCTGGAACACGGCCGCAACTGGTCGGCCATCGCCCGGAT 1920 	1801 GAGCGCCGAGCTGGCCTCCATGGAGCTGAATGAGAGTTCTCGCTGGACAGAAGAAGAAAT 1860 	1741 CCGCATCACCCGCTCAATGGCTAATGAGGCCAACAGCG;GGAGGCCATCACCCCCCAGCA 1800 	81 GAAG 33 GAAG	21 CGAC 73 CGAC	1561 AGATGAGAAGGAGAAAGGAAAGGAGGAGGAGGAGAGAAGCCGGAGGTGGAGAA 1620 	01 GCAC 53 GCAC	GAGO		ωμ

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6697 GGAGCCAGGGCACTCCCGGAGTGCTGTTACCCGCTGCTGTACCGGGATGGGGAACAGAC 6756	6637 GTCGGTCTTGGGTGGTGATGAACAGAACGTATTGAACCTGTGTCCCCACCGGAAGGGCATGAC 6696	6577 CCCGGCCCGTGGCTCCCCCCACAGCGAAGGGGGCAAGAGAGGTCTCCAGAGCCAAACAAGAC 6636	6517 CGTCCTGGACCTCCGCCGCCCACCCAGTGACCTCTACCTCCGCCCCCGGACCATGGTGC 6576	6457 ACAGCAGCTCAGCGCACCCCTGCCCGCCCCCCTCTACTCCTTCCCTGGGGCCAGCTGCCC 6516	6397 GGTCACCCTGGCCCAGCACATCAGTGAGGTCATCACACGGGACTACACCCGGCACCACCC 6456	6337 CCAGCCCTCGTCCAGCCCGCTGCTCCAGACCGCCCCAGGGGTCAAAGGTCAACCAGCGGGT 6396	6277 CCCCGTGAAGCTTGGCGGGGAGGCCGCCCACCTCCCACACCTGCGGCCGCTGCCTGAGAG 6336	6217 GCACCTGGAAGAGCTCGACAAGAGCCACCTGGAGGGGGAGCTGCGGCCCAAGCAGCCAGG 6276	6157 GGTGGAGCCCGTCAGCCCTGTGAGCTCACCCAGTCTGACCCACGACAAGGGGCTCCCCAA 6216	6097 CATCCAGGAACTGGAACTCCGTTCTCTGGGTTACCACGGCAGCAGCTACAGCCCGGAAGG 6156	6037 GGCGCCACCTGCCTCGGCCTCGGACCCGCACCGGGAAAAGACTCAAAGTAAACCCTTTTC 6096	5977 CACCATCGCCCGCACCCCTGCGAAGAACCTCGCACCTCACCACGCCAGCCCGGACCCGGC 6036	5917 CTCCTCCCCAGCAAGGGCTCGGAGCCCCGGCCCCTAGTGCCTCCTGTCTCTGGCCACGC 5976	5857 AGCAGACACCGGCCATGCCTTCCTCGCCAAGCCCCAGCCCGCCTCCGGGCTGGAGCCCGC 5916	5797 CATGGAGCCCGTCTTGCTGCCCAAGGAGGCCCCCGGGTCGCCCGGCCAGAGCGGCCCCG 5856	62 ATTCCC	677 CAGCAAGCCCACGTCCTGAGGT

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Unclassified.

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RS Furness,L.M. and Buchbinder,J.L.

RS Genes expressed in C3A liver cell cultures tro

Patent: US 6673549-A 306 06-JAM-2004;

Incyte Corporation; Palo Alto, CA

Incyte Corporation; Palo Alto, CA

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